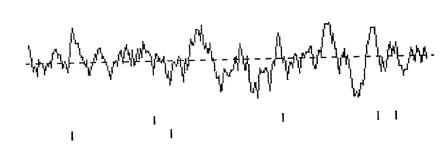
PFAM

Cys Ngly



1 41 81 121 161 201 241 281

>8APro
MLAAPINPSDINMIQGNYGLLPELPAVGGNEGVAQVVAVGSNVTGLKPGDWVIPANAGLG
TWRTEAVFSEEALIQVPSDIPLQSAATLGVNPCTAYRMLMDFEQLQPGDSVIQNASNSGV
GQAVIQIAAALGLRTINVVRDRPDIQKLSDRLKSLGAEHVITEEELRRPEMKNFFKDMPQ
PRLALNCVGGKSSTELLRQLARGGTMVTYGGMAKQPVVASVSLLIFKDLKLRGFWLSQWK
KDHSPDQFKELILTLCDLIRRGQLTAPACSQVPLQDYQSALEASMKPFISSKQILTM



Cys Ngly

1 41 81 121 161 201 241 281 321 361

>8Bpro
MWVCSTLWRVRTPARQWRGLLPASGCHGPAASSYSASAEPARVRALVYGHHGDPAKVVEL
KNLELAAVRGSDVRVKMLAAPINPSDINMIQGNYGLLPELPAVGGNEGVAQVVAVGSNVT
GLKPGDWVIPANAGLGTWRTEAVFSEEALIQVPSDIPLQSAATLGVNPCTAYRMLMDFEQ
LQPGDSVIQNASNSGVGQAVIQIAAALGLRTINVVRDRPDIQKLSDRLKSLGAEHVITEE
ELRRPEMKNFFKDMPQPRLALNCVGGKSSTELLRQLARGGTMVTYGGMAKQPVVASVSLL
IFKDLKLRGFWLSQWKKDHSPDQFKELILTLCDLIRRGQLTAPACSQVPLQDYQSALEAS
MKPFISSKQILTM

adh_zinc: domain 1	of 1, from 1 to 297: score 163.0, E = 5.2e-45 *->pleveevpvpepgpgeVlvkvkaaGiCgsDlhiykgglGlmypgpgd	
8Apro 1	l aa i++sD+ +++g y+ MLAAPINPSDINMIQGNYGLL 2	21
8Apro 22	gthlfpvklPlvlGHEgaGvVeeVGsgVtgfklkvgkfkvGDrVvvlplv + lP v+G Eg+ +V++VGs+Vtg +k+GD V+++ +PELPAVGGNEGVAQVVAVGSNVTGLKPGDWVIPANAG 5	58
	gcCgrgsaeCefCkgsGrenlCpkgratgpgkGlmpndgfggftpkkqga	
8Apro 59	L 5	59
8Apro 60	PckGkdgyhfmgdGgfaEYvvvparrndyfvvkiPdglddeipleeaeaA G++ + +v ++ ++ +P ++ pl AGTWRTEAVFSEEALIQVPSDIPLQSA	35
8Apro 86	allgcaglTaygalvraakvgslppGdtVlVhGaGGGVGlaAvqlAkaaG a lg++ +Tay++l ++++ l+pGd+V + GVG+a++q+A a+G ATLGVNPCTAYRMLMDFEQLQPGDSVIQNASNSGVGQAVIQIAAALG 1	132
8Apro 133	AarViavdssed.pekklelAkelGAdldadfvnnskglptvnddrkedf +++ +v ++d ++ + 1+ +1G a++v+ ++e LRTINVVRDRPDiQKLSDRLK-SLGAEHVITEEELR 1	167
8Apro 168	<pre>veaikeltgGrngagGvDvvldcvGiglggatldaalallkpgGrlvvvG ++k +++ + + +1+c +gg++ ++1+ 1++gG++v+ G RPEMKNFFKDMPQPRLALNCVGGKSSTELLRQLARGGTMVTYG 2</pre>	210
	pkvavGvpgggapipllllkeeeklyersikGsflggrkprlsvlsvd G++ ++ +++ l++k ++++++++++++++++++++++++	
8Apro 211	GMAKQPVVASVsllIFKDLKLRGFWLSQWKKD	242
8Apro 243	ttpdelrealdllasgikdkngkgvldplithtlpplddsleeaneAfel ++pd ++e++ l ++i+ +g+l ++p l + + A+e+ HSPDQFKELILTLCDLIRRGQLTAPACSQVP-LQDYQSALEA	283
8Apro 284	lesgkh.gkvvlip<-* ++ ++ k++l++ SMKPFIsSKQILTM 297	

```
adh zinc: domain 1 of 1, from 57 to 373: score 212.3, E = 7.4e-60
                  *->pleveevpvpepgpgeVlvkvkaaGiCgsDlhiykgglGlmypgpgd
                     ++e++ +++ + +V+vk++aa i++sD+ +++g
                     VVELKNLELAAVRGSDVRVKMLAAPINPSDINMIQGN----YGLL-- 97
      8Bpro
               57
                  gthlfpvklPlvlGHEgaGvVeeVGsgVtgfklkvgkfkvGDrVvvlplv
                        + 1P v+G Eq+ +V++VGs+Vtq
                                                     +k+GD V+++ +
               98 -----PELPAVGGNEGVAQVVAVGSNVTG-----LKPGDWVIPANAG 134
      8Bpro
                  gcCgrgsaeCefCkgsGrenlCpkgratgpgkGlmpndgfggftpkkqga
              135 L----- 135
      8Bpro
                  PckGkdgyhfmgdGgfaEYvvvparrndyfvvkiPdglddeipleeaeaA
                              G++ + +V ++
                                              ++ +P ++
              136 -----GTWRTEAVFSEE----ALIQVPSDI----PLQS---A 161
      8Bpro
                  allgcaglTaygalvraakvgslppGdtVlVhGaGGGVGlaAvqlAkaaG
                  a lq++ +Tay++l ++++ l+pGd+V + GVG+a++q+A a+G
       8Bpro
             162 ATLGVNPCTAYRMLMDFEQ---LQPGDSVIQNASNSGVGQAVIQIAAALG 208
                  AarViavdssed.pekklelAkelGAdldadfvnnskglptvnddrkedf
                   +++ +v ++d ++ + l+ +lG
                                              a++v+
       8Bpro
              209 LRTINVVRDRPDiQKLSDRLK-SLG----AEHVIT------EEELR 243
                  veaikeltgGrngagGvDvvldcvGiglggatldaalallkpgGrlvvvG
                    ++k +++
                               + + +1+c
                                            +gg++ ++l+ l++gG++v+ G
              244 RPEMKNFFKDM---PQPRLALNC----VGGKSSTELLRQLARGGTMVTYG 286
       8Bpro
                  pkvavGvpgggapipl..lllkeeeklyersikGsflgqrkprlsvlsvd
                             ++ +++ l++k
                       G++
                                             ++++G++l+++k +
       8Bpro
              287 ----GMAKQPVVASVslLIFKD-----LKLRGFWLSOWKKD----- 318
                  {\tt ttpdelrealdllasgikdkngkgvldplithtlpplddslee} ane {\tt Afel}
                  ++pd ++e++ l ++i+
                                       +g+l
                                                 ++p 1
       8Bpro
              319 HSPDQFKELILTLCDLIR----RGQLTAPACSQVP-LQ----DYQSALEA 359
                  lesgkh.gkvvlip<-*</pre>
                   ++ ++ k++l++
       8Bpro
              360 SMKPFIsSKQILTM
                                   373
```

FIGURE 4